



## PATENT

## N THE UNITED STATES PATENT AND TRADEMARK OFFICE

APPLICANTS:

SUSANNE LEONHARTSBERGER ET AL. - 3 (PCT)

PCT NO.:

PCT/EP2003/011486

FILED: OCTOBER 16, 2003

SERIAL NO:

10/530,844

FILED: JUNE 5, 2006

TITLE:

FEEDBACK-RESISTANT HOMOSERINE TRANSSUCCINYLASES

WITH A MODIFIED C-TERMINAL

## RESPONSE TO RESTRICTION REQUIREMENT

MAIL STOP AMENDMENT Honorable Commissioner of Patents P.O. Box 1450 Alexandria, VA 22313-1450

Dear Sir:

In response to requirement for restriction dated June 5, 2006, Applicant responds as follows:

Election of Species is on page 2 of this paper
Remarks/Arguments begin on Page 4 of this paper.

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## **ELECTION OF SPECIES:**

The Examiner required a restriction to one of the following four groups for further prosecution:

Group I: Claims 1-9 are drawn to a homoserine transsuccinylase which as compared to wild-type enzyme exhibits a reduced sensitivity to L-methionine or SAM, said polypeptide exhibits a change of at least 2-10 amino acids between the position 285 and 310 of the enzyme as compared with the wild-type enzyme, encoded by polynucleotide with SEQ ID NO: 1 and coding for the corresponding polypeptide of SEQ ID NO:2, vector, host cell (preferably E. coli) and method of making the polypeptide and the method of making L-methionine or SAM.

Group II: Claims 1-9 are drawn to a homoserine transsuccinylase which as compared to wild-type enzyme exhibits a reduced sensitivity to L-methionine or SAM, said polypeptide exhibits a change of at least 2-10 amino acids between the position 285 and 310 of the enzyme as compared with the wild-type enzyme, encoded by polynucleotide with SEQ ID NO: 7 and coding for the corresponding polypeptide of SEQ ID NO:8, vector, host cell (preferably E. coli) and method of making the polypeptide and the method of making L-methionine or SAM.

Group III: Claims 1-9 are drawn to a homoserine transsuccinylase which as compared to wild-type enzyme exhibits a reduced sensitivity to L-methionine or SAM, said polypeptide exhibits a change of at least 2-10 amino acids between the position 285 and 310 of the enzyme as compared with the wild-type enzyme, encoded by polynucleotide with SEQ ID NO:9, and coding for the corresponding polypeptide of SEQ ID NO:10, vector, host cell (preferably E. coli) and method of making the polypeptide and the method of making L-methionine or SAM.

Group IV: Claims 1-9 are drawn to a homoserine transsuccinylase which as compared to wild-type enzyme exhibits a reduced sensitivity to L-methionine or SAM, said polypeptide exhibits a change of at least 2-10 amino acids between the position 285 and 310 of the enzyme as compared with the wild-type enzyme, encoded by polynucleotide with SEQ ID NO: 11 and coding for the corresponding polypeptide of SEQ ID NO: 12, vector, host cell (preferably E. coli) and method of making the polypeptide and the method of making L-methionine or SAM.

**Election**: Applicant elects, with traverse the species of Group II for further prosecution.